RAW SEQUENCE LISTING

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Application Serial Number: _10/593,213_

Source: _1FWP_

Date Processed by STIC: _9/28/06_

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IFWP

RAW SEQUENCE LISTING   DATE: 09/28/2006
PATENT APPLICATION: US/10/593,213   TIME: 11:12:35

Input Set : **A:\2107-299.ST25.txt**
Output Set: **N:\CRF4\09282006\J593213.raw**

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  3  <110> APPLICANT: Lotz, Henrike
  4         Brunner, Herwig
  5         Rupp, Steffen
  7  <120> TITLE OF INVENTION: Hyphae-Specific Cell Wall Proteins of Candida (As Amended)
  9  <130> FILE REFERENCE: P/2107-299
C-->  11  <140> CURRENT APPLICATION NUMBER: US/10/593,213
C-->  12  <141> CURRENT FILING DATE: 2006-09-15
 14  <150> PRIOR APPLICATION NUMBER: 10 2004 013 826.5
 15  <151> PRIOR FILING DATE: 2004-03-16
 17  <160> NUMBER OF SEQ ID NOS: 15
 19  <170> SOFTWARE: PatentIn version 3.3
 21  <210> SEQ ID NO: 1
 22  <211> LENGTH: 336
 23  <212> TYPE: DNA
 24  <213> ORGANISM: Candida Albicans
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 31  acctcagctg cttctggtgc caaatccggt gcttcttcag ttgcttctgc cgctaaatct     180
 33  ggtgtttctt cagctgcctc agctgctaaa tctggtgctt catctgctac cggtggttca     240
 35  tctgctgcca aatctggctc atcaagtggt gccggttttg ctcctgtcgc tggtgctggt     300
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 52              20                  25                  30
 55  Ala Ser Gly Ala Lys Ser Gly Ala Thr Ser Ala Ala Ser Gly Ala Lys
 56          35                  40                  45
 59  Ser Gly Ala Ser Ser Val Ala Ser Ala Ala Lys Ser Gly Val Ser Ser
 60      50                  55                  60
 63  Ala Ala Ser Ala Ala Lys Ser Gly Ala Ser Ser Ala Thr Gly Gly Ser
 64  65                  70                  75                  80
 67  Ser Ala Ala Lys Ser Gly Ser Ser Ser Gly Ala Gly Phe Ala Pro Val
 68              85                  90                  95
 71  Ala Gly Ala Gly Ser Leu Ala Ala Ile Ala Gly Leu Leu Leu Leu
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85  aagaccgatt atattaaatt ttttgccacc gcaaaagatg ttccaggtga tttatctacg     180
87  ttggctacca aagtgttgac ttatactgat gattcataca caactttgtt gaatgatgat     240
89  tctttgaatg tttccaactt agaagcatat gctactagtt tgccatggta ttccagaatt     300
91  caagctgatg ctggtggcaa aggttctgcc tccggttctg cctctggctc tggttctgcc     360
93  aaatcaactg caagtgctga aaaatctagt ggctcaagtg cttctgcttc aagcactgca     420
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112             20                  25                  30
115 Leu Val Gly Asp Tyr Gln Asp His Lys Thr Asp Tyr Ile Lys Phe Phe
116         35                  40                  45
119 Ala Thr Ala Lys Asp Val Pro Gly Asp Leu Ser Thr Leu Ala Thr Lys
120     50                  55                  60
123 Val Leu Thr Tyr Thr Asp Asp Ser Tyr Thr Thr Leu Leu Asn Asp Asp
124 65                  70                  75                  80
127 Ser Leu Asn Val Ser Asn Leu Glu Ala Tyr Ala Thr Ser Leu Pro Trp
128                 85                  90                  95
131 Tyr Ser Arg Ile Gln Ala Asp Ala Gly Gly Lys Gly Ser Ala Ser Gly
132             100                 105                 110
135 Ser Ala Ser Gly Ser Gly Ser Ala Lys Ser Thr Ala Ser Ala Glu Lys
136         115                 120                 125
139 Ser Ser Gly Ser Ser Ala Ser Ala Ser Ser Thr Ala Gly Gly Ser Ser
140     130                 135                 140
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152 <211> LENGTH: 1682
153 <212> TYPE: DNA
154 <213> ORGANISM: Candida Albicans
156 <400> SEQUENCE: 5
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159 cttgttatca ctcaaaacag agtcgatcgt ggtgttcttg acgttagtgt tggaaatatc     120
161 accatcaatt ctggagcttc ttggtcaatt atcaacaatg ctatatcaac ccttgttgga     180
163 agtttaactg ttcagcccaa tgctggtctt tacattaccc tgacttcacc ccttttgtca     240
165 cttcaagtca cattaacttc tttgcttagc acaattcaaa acaatggtat tattgcgttc     300
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171 gcatcttggt caaacagtgg attgatggca ttttatcaaa atcaaagaag ttctggtatt   480
173 gttagtcttg ggacaccatc aggttcaata accaataatg gtcaaatctg tttgattaac   540
175 gaagtctaca aacagaccac aagcatcaac ggttctggtt gtttcactgc caatcgtaac   600
177 tcgacaatat atattgccaa tgtattgtta ccagtttcca catcgcaaaa ttttttatttg   660
179 gcagacagcc aatcttccat aattgttcaa gctatttcaa cccctcaagt gtttaatgtc   720
181 tatgggtttg gtaacggtaa tatggtcggg gttactcttc cattgatcgg taatatatgg   780
183 aatccagcat atagttataa tccatccaca ggtattttaa gattgagaaa tttttttgtg   840
185 tatcaagatt ttaatattgg tcctggttat aatcctagtt tatttctgat cgttactgac   900
187 aatggtgctg gtcttccctc aacaatactc ggttcggttt cttatagtgg tcctgttcca   960
189 ccaagagctt tacccgcatc ttgtaagatt gcatgtaaac ccgtgcctac cgcgccagga  1020
191 actaatccaa ccgagtacac gaccacaata acaacaacaa attctgctgg taagccattg  1080
193 acagaaactg gtgtggttga tattctgact gatagtaacg gatcatggtt ctcaagtact  1140
195 acaatctttc caacttcgtc gtcaagtagt agcagtagca gcactgtttc ttcaactgct  1200
197 ccgtcatcct caagcaccaa accttcatcc agtagccaac catcttctac tccaccacca  1260
199 tcttcaagta gtaaagcatc atcaactact ccaagctcta gtagtcaatc gtcttcaact  1320
201 actccaagct caagcagtaa gccttcctca actgtaccac caactggcag cagtcagtca  1380
203 tcttcaacta tcccaagttc cagtactcaa ccttcttcta ctgctccatc atctttaagt  1440
205 tctccatctt cttctactac tccaagctcc agcagtcaat cttcattttc tgctcaaagc  1500
207 tctattggcc agacatcgtc ttctactgta tcttcgtcga gtagtcaacc atcgtgctgg  1560
209 gagtcatcaa gcagtcagtc gtcatccggt acgacaagtt ccagtagtca gttttcttca  1620
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213 tc                                                                 1682
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221 <400> SEQUENCE: 6
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228             20                  25                  30
231 Leu Asp Val Ser Val Gly Asn Ile Thr Ile Asn Ser Gly Ala Ser Trp
232         35                  40                  45
235 Ser Ile Ile Asn Asn Ala Ile Ser Thr Leu Val Gly Ser Leu Thr Val
236     50                  55                  60
239 Gln Pro Asn Ala Gly Leu Tyr Ile Thr Ser Thr Ser Pro Leu Leu Ser
240 65                  70                  75                  80
243 Leu Gln Val Thr Leu Thr Ser Leu Leu Ser Thr Ile Gln Asn Asn Gly
244             85                  90                  95
247 Ile Ile Ala Phe Asn Ser Ser Pro Ser Leu Thr Ser Ser Thr Tyr Asn
248             100                 105                 110
251 Leu Val Gly Leu Ser Leu Val Asn Thr Gly Glu Met Tyr Phe Ser Ala
252         115                 120                 125
255 Ser Gly Val Leu Pro Ser Val Met Ala Leu Thr Ala Ala Ser Trp Ser
256     130                 135                 140
259 Asn Ser Gly Leu Met Ala Phe Tyr Gln Asn Gln Arg Ser Ser Gly Ile
260 145                 150                 155                 160
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263 Val Ser Leu Gly Thr Pro Ser Gly Ser Ile Thr Asn Asn Gly Gln Ile
264               165                 170                 175
267 Cys Leu Ile Asn Glu Val Tyr Lys Gln Thr Thr Ser Ile Asn Gly Ser
268               180                 185                 190
271 Gly Cys Phe Thr Ala Asn Arg Asn Ser Thr Ile Tyr Ile Ala Asn Val
272           195                 200                 205
275 Leu Leu Pro Val Ser Thr Ser Gln Asn Phe Tyr Leu Ala Asp Ser Gln
276       210                 215                 220
279 Ser Ser Ile Ile Val Gln Ala Ile Ser Thr Pro Gln Val Phe Asn Val
280 225                 230                 235                 240
283 Tyr Gly Phe Gly Asn Gly Asn Met Val Gly Val Thr Leu Pro Leu Ile
284               245                 250                 255
287 Gly Asn Ile Trp Asn Pro Ala Tyr Ser Tyr Asn Pro Ser Thr Gly Ile
288               260                 265                 270
291 Leu Arg Leu Arg Asn Phe Phe Val Tyr Gln Asp Phe Asn Ile Gly Pro
292           275                 280                 285
295 Gly Tyr Asn Pro Ser Leu Phe Ser Ile Val Thr Asp Asn Gly Ala Gly
296       290                 295                 300
299 Leu Pro Ser Thr Ile Leu Gly Ser Val Ser Tyr Ser Gly Pro Val Pro
300 305                 310                 315                 320
303 Pro Arg Ala Leu Pro Ala Ser Cys Lys Ile Ala Cys Lys Pro Val Pro
304               325                 330                 335
307 Thr Ala Pro Gly Thr Asn Pro Thr Glu Tyr Thr Thr Thr Ile Thr Thr
308               340                 345                 350
311 Thr Asn Ser Ala Gly Lys Pro Leu Thr Glu Thr Gly Val Val Asp Ile
312           355                 360                 365
315 Ser Thr Asp Ser Asn Gly Ser Trp Phe Ser Ser Thr Thr Ile Phe Pro
316       370                 375                 380
319 Thr Ser Ser Ser Ser Ser Ser Ser Ser Ser Thr Val Ser Ser Thr Ala
320 385                 390                 395                 400
323 Pro Ser Ser Ser Ser Thr Lys Pro Ser Ser Ser Ser Gln Pro Ser Ser
324               405                 410                 415
327 Thr Pro Pro Pro Ser Ser Ser Ser Lys Ala Ser Ser Thr Thr Pro Ser
328               420                 425                 430
331 Ser Ser Ser Gln Ser Ser Ser Thr Thr Pro Ser Ser Ser Ser Lys Pro
332           435                 440                 445
335 Ser Ser Thr Val Pro Pro Thr Gly Ser Ser Gln Ser Ser Ser Thr Ile
336       450                 455                 460
339 Pro Ser Ser Ser Thr Gln Pro Ser Ser Thr Ala Pro Ser Ser Leu Ser
340 465                 470                 475                 480
343 Ser Pro Ser Ser Ser Thr Thr Pro Ser Ser Ser Ser Gln Ser Ser Phe
344               485                 490                 495
347 Ser Ala Gln Ser Ser Ile Gly Gln Thr Ser Ser Ser Thr Val Ser Ser
348           500                 505                 510
351 Ser Ser Ser Gln Pro Ser Cys Trp Glu Ser Ser Ser Ser Gln Ser Ser
352       515                 520                 525
355 Ser Gly Thr Thr Ser Ser Ser Ser Gln Phe Ser Ser Ser Ala Pro Pro
356       530                 535                 540
359 Ser Ser Thr Gln Ser Ser Phe Thr Ala Glu Ser Ser Asn Ser Gln Leu
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Input Set : A:\2107-299.ST25.txt
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364 <211> LENGTH: 30
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373 <211> LENGTH: 29
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375 <213> ORGANISM: Candida Albicans
377 <400> SEQUENCE: 8
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382 <211> LENGTH: 36
383 <212> TYPE: DNA
384 <213> ORGANISM: Candida Albicans
386 <400> SEQUENCE: 9
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391 <211> LENGTH: 27
392 <212> TYPE: DNA
393 <213> ORGANISM: Candida Albicans
395 <400> SEQUENCE: 10
396 cgagctcccg aaatgccacc atagttt                                 27
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400 <211> LENGTH: 29
401 <212> TYPE: DNA
402 <213> ORGANISM: Candida Albicans
404 <400> SEQUENCE: 11
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409 <211> LENGTH: 29
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411 <213> ORGANISM: Candida Albicans
413 <400> SEQUENCE: 12
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418 <211> LENGTH: 36
419 <212> TYPE: DNA
420 <213> ORGANISM: Candida Albicans
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VERIFICATION SUMMARY   DATE: 09/28/2006
PATENT APPLICATION: US/10/593,213   TIME: 11:12:36

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L:11 M:270 C: Current Application Number differs, Replaced Current Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date